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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO
10/712,425	11/13/2003	Frank D. Lee	EPT-001C1	9956
51414	7590 07/28/2006		EXAMINER	
GOODWIN PROCTER LLP			LIN, JERRY	
PATENT AL EXCHANGE	MINISTRATOR E PLACE		ART UNIT PAPER NUMBER	
BOSTON, N	1A 02109-2881		1631	
			DATE MAILED: 07/28/2006	

Please find below and/or attached an Office communication concerning this application or proceeding.

		Application No.	Applicant(s)			
Office Action Summary						
		10/712,425 Examiner	LEE ET AL.			
	·		Art Unit			
The MAILING	G DATE of this communication app	Jerry Lin ears on the cover sheet with the c	1631			
Period for Reply			orrospondente address =			
WHICHEVER IS LC - Extensions of time may be after SIX (6) MONTHS fir - If NO period for reply is s - Failure to reply within the Any reply received by the	ATUTORY PERIOD FOR REPLY DNGER, FROM THE MAILING DA e available under the provisions of 37 CFR 1.13 om the mailing date of this communication. pecified above, the maximum statutory period w set or extended period for reply will, by statute, office later than three months after the mailing tment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 6(a). In no event, however, may a reply be tim iiii apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	I. lely filed the mailing date of this communication.			
Status						
1) Responsive to	communication(s) filed on 09 Ju	ne 2006.				
, <del></del>	This action is <b>FINAL</b> . 2b)⊠ This action is non-final.					
	Since this application is in condition for allowance except for formal matters, prosecution as to the merits is					
closed in accordance with the practice under Ex parte Quayle, 1935 C.D. 11, 453 O.G. 213.						
Disposition of Claims						
4a) Of the abo 5) ☐ Claim(s) 6) ☑ Claim(s) <u>1-10</u> 7) ☐ Claim(s)	and 126-132 is/are pending in the eve claim(s) 11,15,26-30 and 128 in is/are allowed.  12-14,16-25,31-35,126,127 and in is/are objected to.  12-14 are subject to restriction and/or	is/are withdrawn from considerat 129-132 is/are rejected.	ion.			
Application Papers						
10) ☐ The drawing(s Applicant may i Replacement d	on is objected to by the Examiner ) filed on is/are: a) acce not request that any objection to the d rawing sheet(s) including the correction coloration is objected to by the Examine	pted or b) objected to by the E rawing(s) be held in abeyance. See on is required if the drawing(s) is obj	37 CFR 1.85(a). ected to. See 37 CFR 1.121(d).			
Priority under 35 U.S.(	C. § 119					
<ul> <li>12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).</li> <li>a) All b) Some * c) None of:</li> <li>1. Certified copies of the priority documents have been received.</li> <li>2. Certified copies of the priority documents have been received in Application No.</li> <li>3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).</li> <li>* See the attached detailed Office action for a list of the certified copies not received.</li> </ul>						
	s Patent Drawing Review (PTO-948) Statement(s) (PTO-1449 or PTO/SB/08)	4) Interview Summary ( Paper No(s)/Mail Dai 5) Notice of Informal Pa 6) Other:				

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# **DETAILED ACTION**

#### Election/Restrictions

1. Applicant's election with traverse of Group I in the reply filed on March 15, 2006 is acknowledged. The traversal is on the ground(s) that Groups I-IV share common technical features and the search of Groups I-IV would not be an undue burden on the Examiner. This is not found persuasive because Groups II-IV all contain different features from Group I that would require a separate search. Although the Groups do share common technical features, the Groups also contain features that differentiate themselves from the other groups, as was explained in the previous office action. A thorough search of the art would require the Examiner to also search for these different features. For example, Group II has the limitations of arrays, solid supports or a surface plasmon resonance array, none of which are found in Group I. Since the each of these additional features would require separate searches for each group, a search for Groups I-IV would be an undue search burden on the Examiner.

The requirement is still deemed proper and is therefore made FINAL.

2. Furthermore, in a Supplemental Species Election, Applicants elected without traverse of phosphorylation as the post translational modification and tyrosine as the location of post-translational modification in Species A; full length antibody as the type of capture agent in Species B; claim 14 in species C; and fluorescent labels as a type of secondary capture agent in Species D, in the reply filed on June 9, 2006.

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#### Status of the Claims

Claims 1-10, 12-14, 16-25, 31-35, 126, 127, and 129-132 are under examination.

Claims 11, 15, 26-30, 128 are withdrawn as being directed toward an unelected invention. Instant claim 128 is drawn to a stainable dye, not a fluorescent label.

Claims 36-125 are cancelled (claims 37-125 are drawn to an unelected invention).

## Specification

3. The disclosure is objected to because it contains an embedded hyperlink and/or other form of browser-executable code (for example, page 118, line 2 or page 121, line 18). Applicant is required to delete the embedded hyperlink and/or other form of browser-executable code. See MPEP § 608.01.

# Claim Rejections - 35 USC § 112, 2<sup>nd</sup> Paragraph

4. The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 1-10, 12-14, 16-25, 31-35, 126, 127, and 129-132 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

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Claim 1 recites the limitation "said treatment" in line 11. There is insufficient antecedent basis for this limitation in the claim.

Claim 18 is unclear because of the presence of the parenthetical terms. It is unclear if the terms in the parentheses are intended to limit the claim or if the other terms are intended to limit the claim.

Claim 19 is also unclear because it is unclear if the term "treatment" is intended to refer to the "treatment" in claim 1 or some other treatment.

## Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35U.S.C. 102 that form the basis for the rejections under this section made in thisOffice action:

A person shall be entitled to a patent unless -

(e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351(a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

Claims 1-3, 7-10, 12-14, 16-22, 31- 35, 126, and 129 are rejected under 35 U.S.C. 102(e) as being anticipated by Katz (US 2002/0137119 A1).

6. The instant claims are drawn to method of detecting the presence of post-translational modification on a sample that include the steps of identifying potential post-translational modification sites and a proteome epitope tag on the fragments of a protein from a sample, generating a capture agent that binds to the proteome epitope tag, subjecting the sample to a treatment to solublize the

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fragments, and detecting the presence or absence of a post-translational modification. It is noted that the term "Proteome Epitope Tag" is not specifically defined in the specification, thus this phrase has been interpreted to mean epitopes that identify a protein.

Regarding claim 1, Katz teaches a method that computationally identifies amino acid sequences from a sample of proteins wherein each fragment contains a unique proteome epitope tag (i.e., unique antigen) (page 2, paragraphs 0020-0022; page 4, paragraph 0062; page 5, paragraph 0082) and a post-translational modification site (page 5, paragraph 0076; page 6, paragraphs 0091, 0093); generating a capture agents that bind to the PET (page 4, paragraph 0068) which may be immobilized to a support (page 8, paragraph 0140); subjecting the sample to render the fragment soluble in solution (page 8, paragraph 0137-0139); detecting the presence or absence of post-translational modification (page 6, paragraph 0091).

Regarding claims 2, 3 and 126, Katz teaches wherein the post-translational modification is phosphorylation is on tyrosine (page 6, paragraph 0091).

Regarding claim 7, Katz teaches analyzing amino acid sequences in terms of solubility (page 5, paragraph 0076).

Regarding claim 8, Katz teaches that the fragments are 5-12 amino acids in length (page 6, paragraph 0094).

Regarding claim 9, 10 and 129, Katz teaches that the capture agents may be full-length antibodies (proteins) (page 7, paragraphs 0126-0129).

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Regarding claims 12 and 16, Katz teaches wherein the treatment is by a protease or chemical agent (page 3, paragraphs 0043-0044; page 8, paragraph 0138).

Regarding claims 13 and 14, Katz teaches denaturizing using chemical or thermo means (page 12, paragraph 0171; page 13, paragraph 0178).

Regarding claims 17 and 18, Katz teaches that the sample may be obtained from a variety of sources (page 8, paragraph 0136; page 7, paragraph 0125; page 14, paragraph 0187).

Regarding claim 19, Katz teaches that a sample is produced by treatment of membrane bound proteins (page 9, paragraph 0157).

Regarding claims 20 and 21, Katz teaches where the post-translational modification is preserved and does not overlap with the PET (page 6, paragraph 0091-0093).

Regarding claim 22, Katz et al. teach optimizing the specificity of antibodies (page 4, paragraph 0058; page 7, paragraph 121-123, 130).

Regarding claim 31, Katz et al. teach identifying a fragment from a sample containing unrelated proteins (page 14, paragraph 0187-189).

Regarding claim 32, Katz teaches determining the amount of fragments bound to capture agents (page 8, paragraph 0143).

Regarding claim 33, Katz teaches where capture agents are produced by immunizing animals with the antigen with the PET sequence (page 7, paragraph 0125).

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Regarding claim 34 and 35, Katz teaches labeling the protein fragments which would block the N or C-terminus of the PET sequence by a chemical group (page 13, paragraph 0175).

#### Claim Rejections - 35 USC § 103

- 7. The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:
  - (a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.

This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(e), (f) or (g) prior art under 35 U.S.C. 103(a).

8. Claims 4-6 are rejected under 35 U.S.C. 103(a) as being unpatentable over Katz (US 2002/137119 A1) in view of Whaley et al. (Biological Mass Spectrometry, (1991) Volume 20, pages 210-214).

The instant claims are drawn to a method of computationally creating fragments from a sample of proteins and identifying unique proteome epitope tags with a Nearest Neighbor analysis, where each fragment contains a unique proteome epitope tag and obtaining capture agents that selectively binds to the proteome epitope tags.

Katz is applied as above.

Although Katz teaches computationally analyzing the plurality of fragments according to one parameter defining characteristic of an amino acid sequence (page 2, paragraph 0021) and identifying unique proteome tags (antigens), Katz does not explicitly teach where the parameter is a Nearest Neighbor Analysis that identifies proteome epitope tags.

Regarding claim 4, Whaley et al. teach a method wherein the protein digests (fragments) are analyzed using Nearest Neighbor analysis based on charge (page 212, left column).

It would have been obvious to one of ordinary skill in the art at the time of the invention to combine the methods of Katz with Whaley et al. to gain the advantage of ranking proteins sequence in order of their importance. Katz discloses that it is desirable to rank the proteins in accordance to the importance of parameters (page 5, paragraph 0078) to allow the user to identify which sequences have the most desirable traits. However, Katz does not teach a specific method of ranking the proteins sequences. Thus one of ordinary skill in the art would be motivated to find such a ranking system to use in Katz's methods. Nearest Neighbor analysis is a well-known method of classifying

objects in accordance to the user's desired characteristics and with high confidence. Whaley et al. teach applying the Nearest Neighbor analysis to protein fragments. Thus one seeking to use a ranking system for identifying a desired fragment, would seek to use the Nearest Neighbor analysis since it may be applied to protein fragments, as demonstrated by Whaley et al, and since the analysis produces results with a high degree of confidence. Thus it would have been obvious to one of ordinary skill in the art to combine the methods of Katz with Whaley et al.

Regarding claims 5 and 6, Katz teaches that the capture agents are determined to have the desired high specificity with minimal cross-reactivity (page 4, paragraph 0058; page 7, paragraph 121-123, 130).

9. Claims 23-25, 127, and 130-132 are rejected under 35 U.S.C. 103(a) as being unpatentable over Katz (US 2002/137119 A1) in view of Gembitsky et al. (US 2005/0153298).

The instant claims are drawn to method of detecting the presence of post-translational modification on a sample that include the steps of identifying potential post-translational modification sites and a proteome epitope tag on the fragments of a protein from a sample, generating a capture agent that binds to the proteome epitope tag, subjecting the sample to a treatment to solublize the fragments, and detecting the presence or absence of a post-translational

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modification. In addition, the instant claims also include steps of using a secondary capture agent specific for post-translational modification.

Katz is applied as above.

Although Katz teaches identifying post-translational modifications (page 5, paragraph 0076; page 6, paragraphs 0091, 0093), Katz does not specifically teach using a secondary capture agent.

Regarding claims 23-25, 127, and 130-132, Gembitsky et al. teach using a fluorescent-labeled secondary capture agent (full-length antibody) specific for phosphorylated tyrosine (page 3, paragraphs 0033, 0034; page 4, paragraphs 0039, 0040; page 5, paragraph 0046; page 7, paragraph 0074).

It would have been obvious to one of ordinary skill in the art at the time of the invention to combine the methods of Katz with Gembitsky et al. to gain the advantage of high throughput analysis of cellular protein modifications. Katz teaches creating protein fragments and then binding the fragments to immobilized antibodies (page 8, paragraph 0140). Katz also teaches detecting the presence or absence of post-translational modifications (page 5, paragraph 0076; page 6, paragraphs 0091, 0093). Katz teaches using well-known techniques for detecting post-translational modifications. However Gembitsky et al. teach that these well-known techniques lack expression profiling that allows parallel quantitation of all proteins expressed in a cell or tissue that analyzes the type, degree and timing of dynamic post-translational protein modifications (page 2, paragraph 0013). Gembitsky et al. teach that their method fulfills this need and allows the practitioner to perform a high throughput and quantitative method

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of comparative analysis of post-translational protein modifications (page 2, paragraph 0016). Thus one of ordinary skill in the art would be motivated to combine the methods of Katz with Gembitsky et al. to gain the advantages taught by Gembitsky et al.

#### Contact Information

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Jerry Lin whose telephone number is (571) 272-2561. The examiner can normally be reached on 10:00am-6:30pm M-F.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Andrew Wang, can be reached on (571) 272-0811. The fax phone number for the organization where this application or proceeding is assigned is (571) 273-8300.

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JL

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